

Fig. 1

a0022 (LIP9)

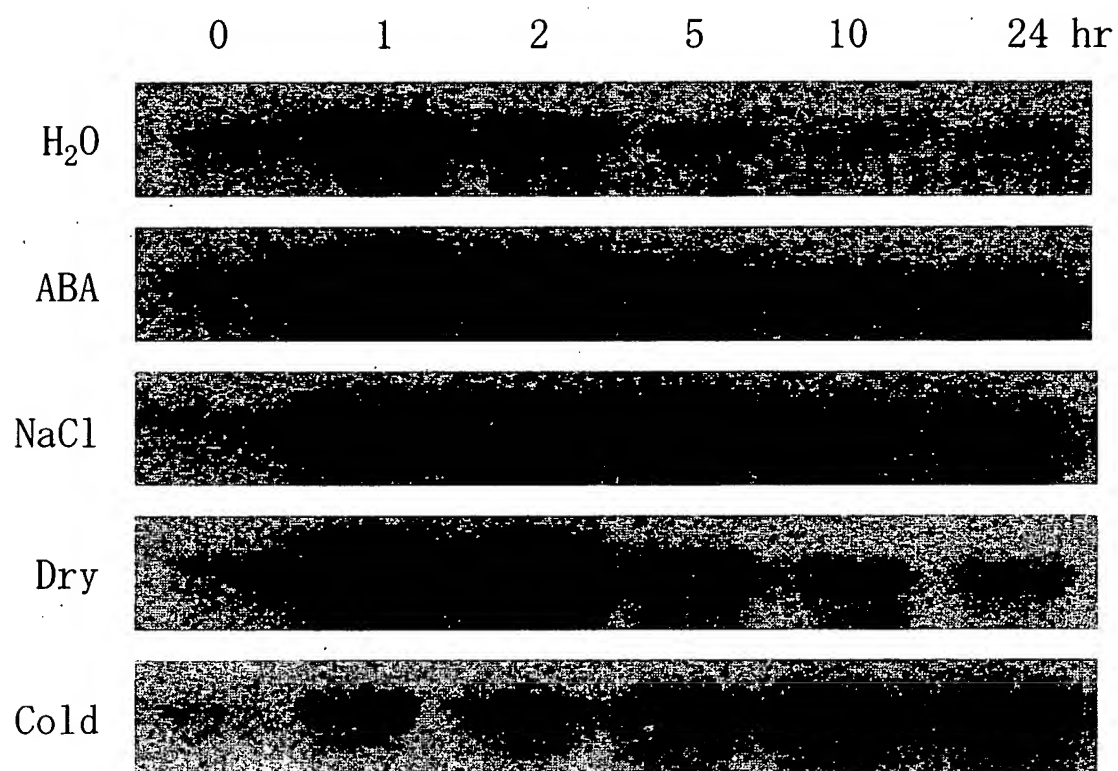


Fig. 1

a0022 (LIP9)

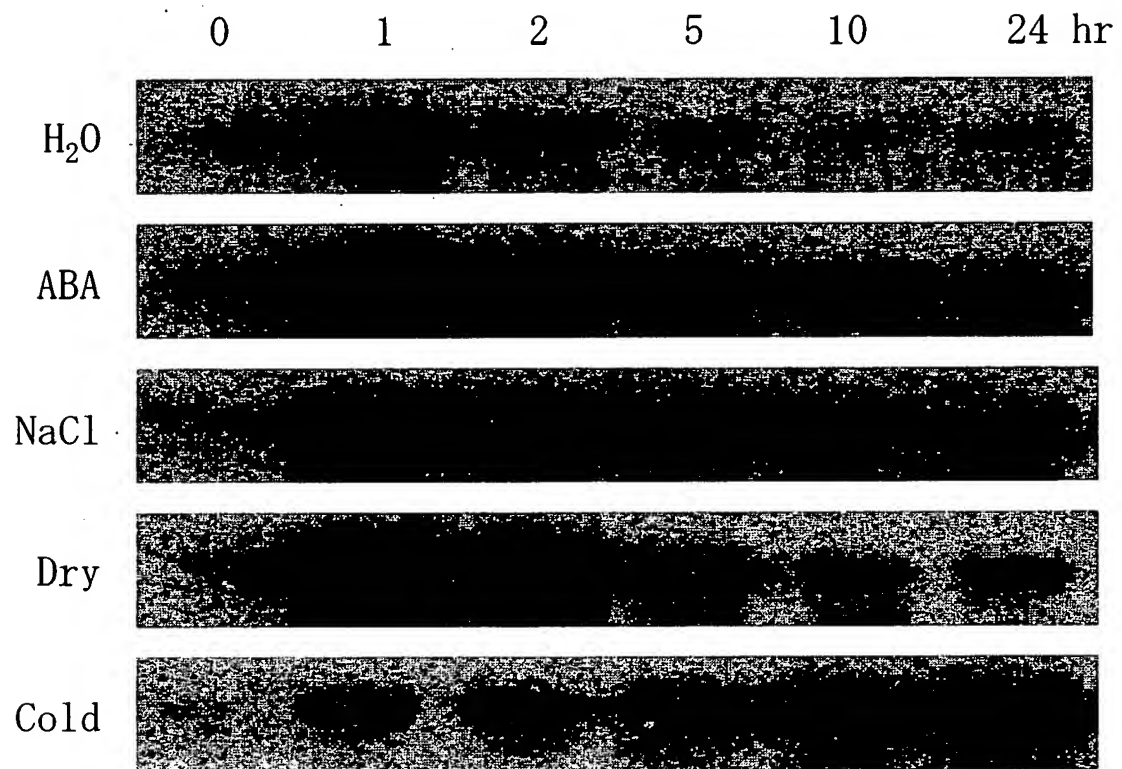


Fig. 1

a0022 (LIP9)

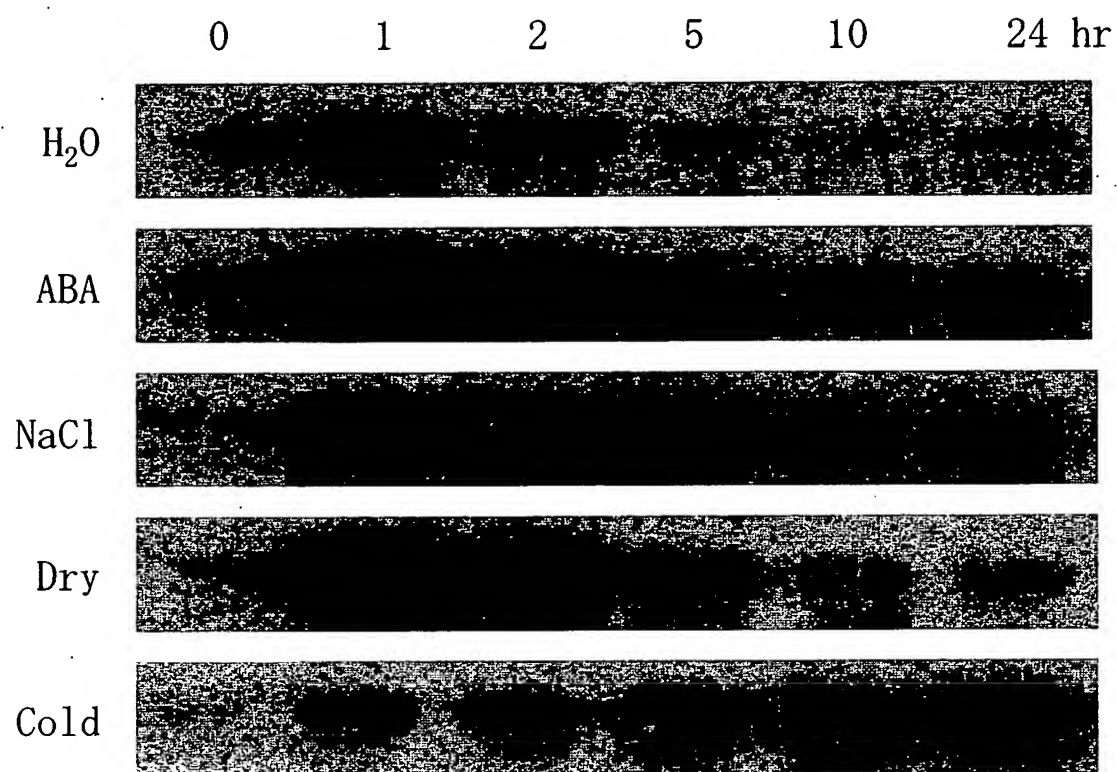


Fig. 2

## LIP 9

10	20	30	40	50	60
TCATCAGCTA	TCATCAAAGC	GAAGGAAAGA	AAGAAAAATA	AAAGGAAAAG	AACTGGCTGG
70	80	90	100	110	120
AAATTAGAGA	AGCCCCGGAC	GACTCGATCT	GGGGGTGGCA	AATTAATCAG	TGTGATCAAC
130	140	150	160	170	180
AGGGGATAACT	TATCCCGTCC	GACCAAATCC	ACCAACCAAA	CCAAGACCCG	ATTTGTTAGG
190	200	210	220	230	240
CTGTGAAAGA	CGGATCAGTG	GGACCCTGAT	CTACGGACCC	CATATGTCAC	CGTCCAGGTC
250	260	270	280	290	300
TCTGGATCTC	TCCCGTCTGC	CTAATCAGAC	ACCGCGCGCG	CGGTGCCGTC	GCTCTCGAGC
310	320	330	340	350	360
CGTGTCCCGC	TCCCAACTCG	TCACAAAAGC	GATCACAGAC	TCTTCCTTCC	TCTGCTGGGA
370	380	390	400	410	420
GAGAAGAAAA	ATTGGCCGCG	ATGATGCCGA	TAAAGAGGAA	AAAGGGATGA	GAATCCGATG
430	440	450	460	470	480
GAAAAAAACT	GATGTTAATC	TATCGCTACT	GCTGCGCACT	AAGACGAATC	GTATCCGAAC
490	500	510	520	530	540
AAGAAACGCT	TACGTTACTG	TTCCTAAATG	GATCGTCCG	CTCATCACTT	AACCAAAAAAT
550	560	570	580	590	600
CGATTAGGAA	ATTGACGGAC	AGCGACGCC	GAAGCCAAGT	GTCTCGTCGC	GTAGGGCTCG
610	620	630	640	650	660
AGGCCTCGAA	GCAGAGGGAG	CGGAGAGGCG	GACGCGCCG	CCACGCCTCC	TCTCCCTCGG
670	680	690	700	710	720
TGACACGGCC	GTCTGGCTCC	ACATGGCGCC	GACCTCTCCC	GATGCGTCCA	CCCGTCCCGA
730	740	750	760	770	780
GGCACCGCCA	CGTCGGAACC	AGCCGGCCGC	CCCACGCGAT	TGCCGACACG	CGTCGCGGCG
790	800	810	820	830	840
CCACTGGCTC	ACCCGCTGCC	TGCCTCTGCC	TGCCCCCAT	CTCGTCGCCA	TTTCCCGCCC
850	860	870	880	890	900
ACGCTTCTTG	TCCTCGCGTC	GCCTACGCGT	ACGTACGATA	CAAACGCCGC	ACCTTTCGAT
910	920	930	940	950	960
CCCCTCCGCT	ATATAAGGAG	GGCATCTGCC	TCGCCACCTT	CTTCATCCGA	AAGCAAAAGC
970	980	990	1000	1010	1020
GACTCGTCAC	AGCTCAAACA	AGTCAAGAGC	GAATAGTTCT	TGCTGATCTG	TTGTTTGATT
1030	1040	1050	1060	1070	1080
ACTTTAGTTC	TCGAGAGGCT	TTAGCTGAAT	CCATCGATCA	TGGAGGATGA	GAGGAACA..

**LIP 9**

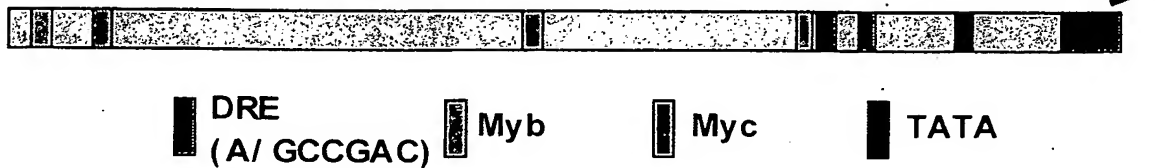


Fig. 3

LIP9: Gus construct

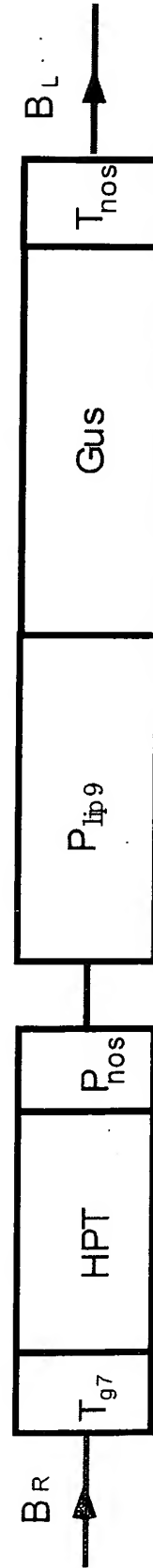


Fig. 4

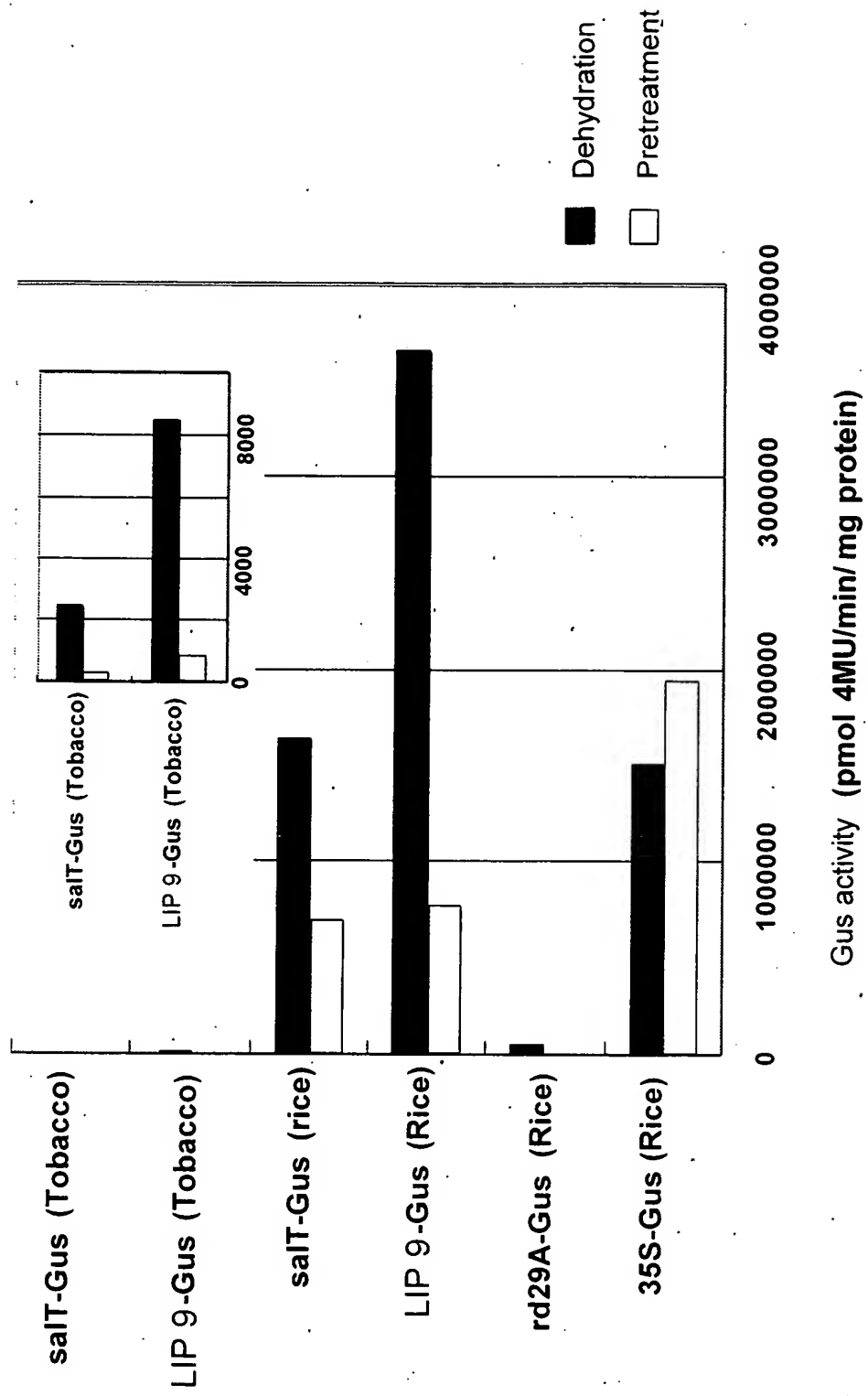
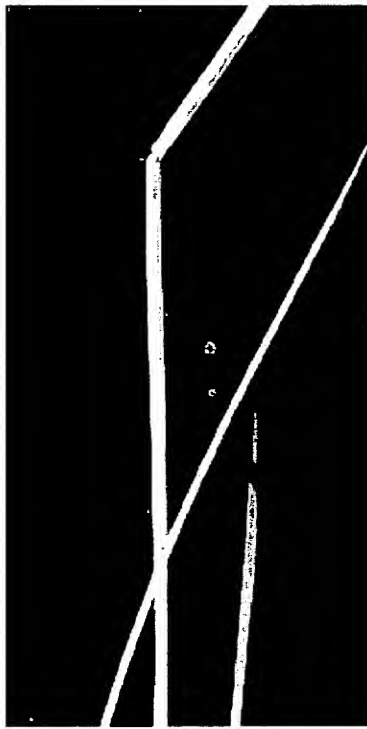
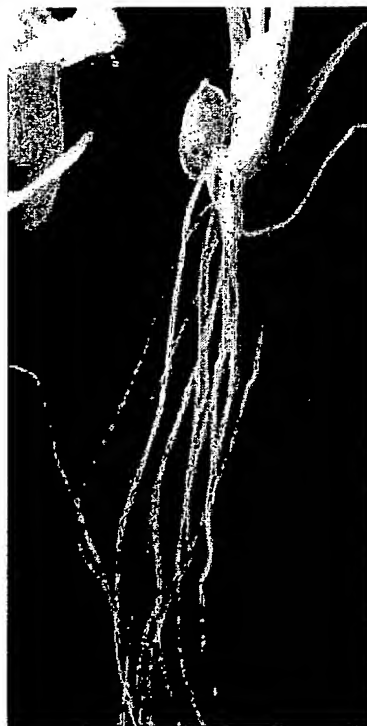


Fig. 5

Leaves



Roots

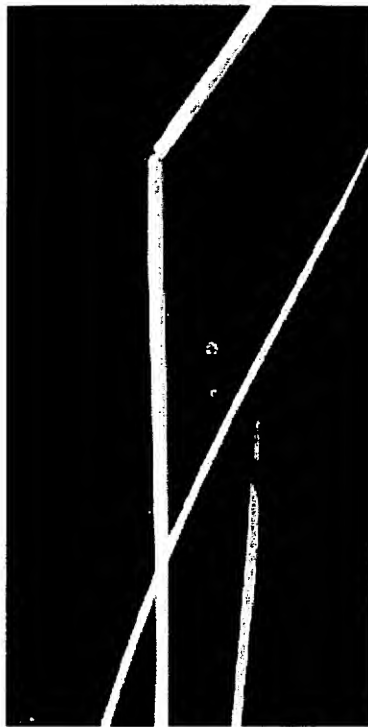


control

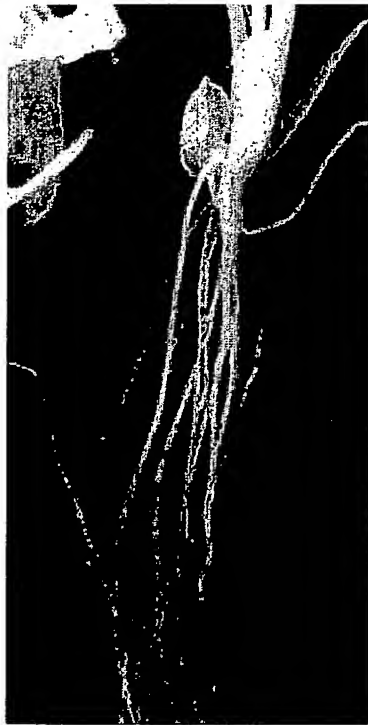
Salt stress

Fig. 5

Leaves



Roots

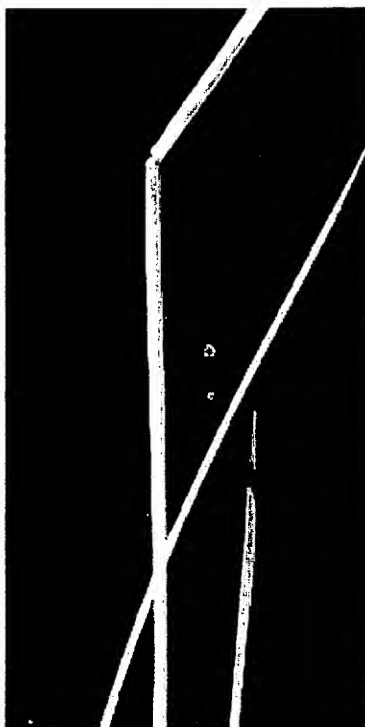


control

Salt stress

Fig. 5

Leaves



Roots



control

Salt stress

Fig. 6

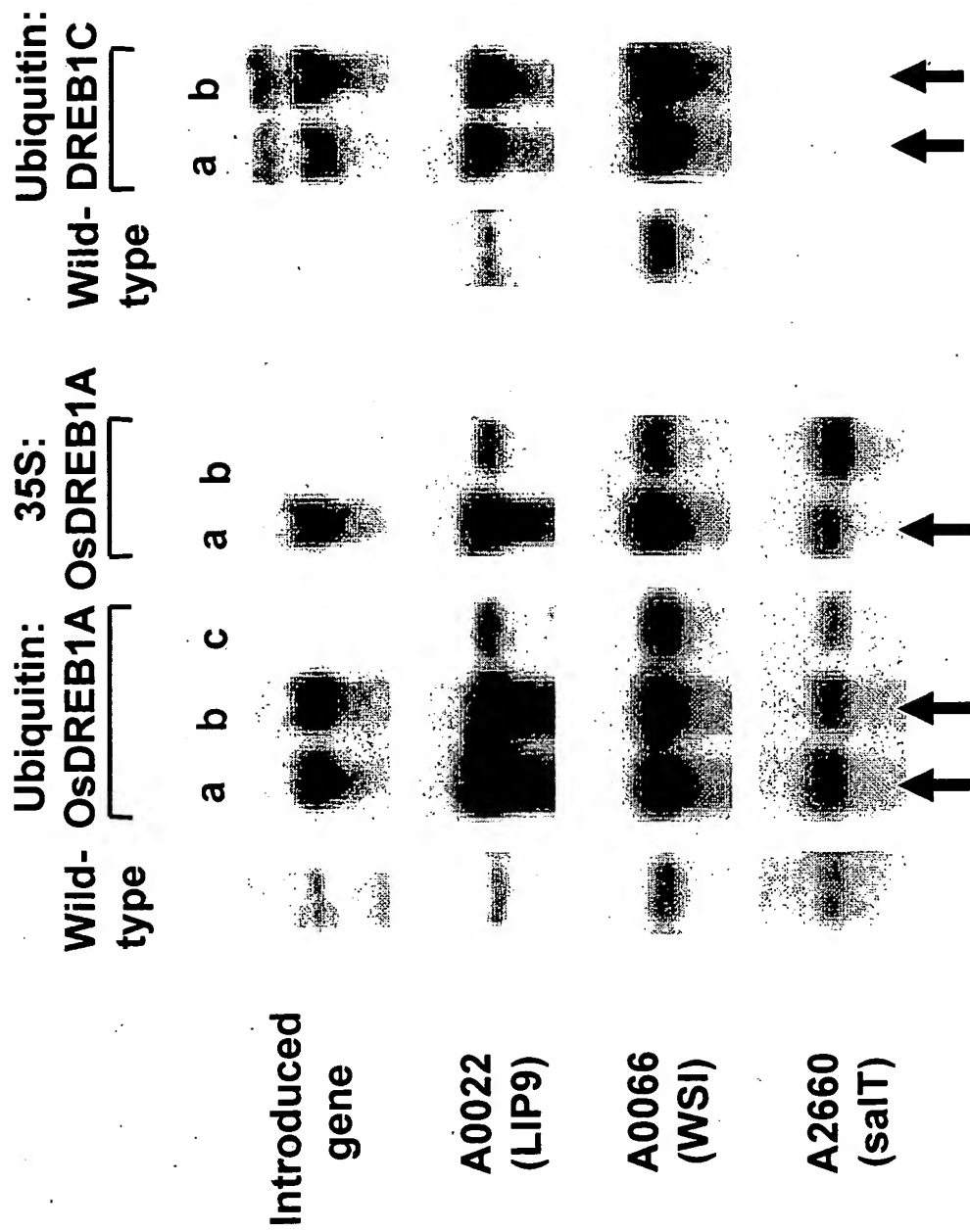


Fig. 6

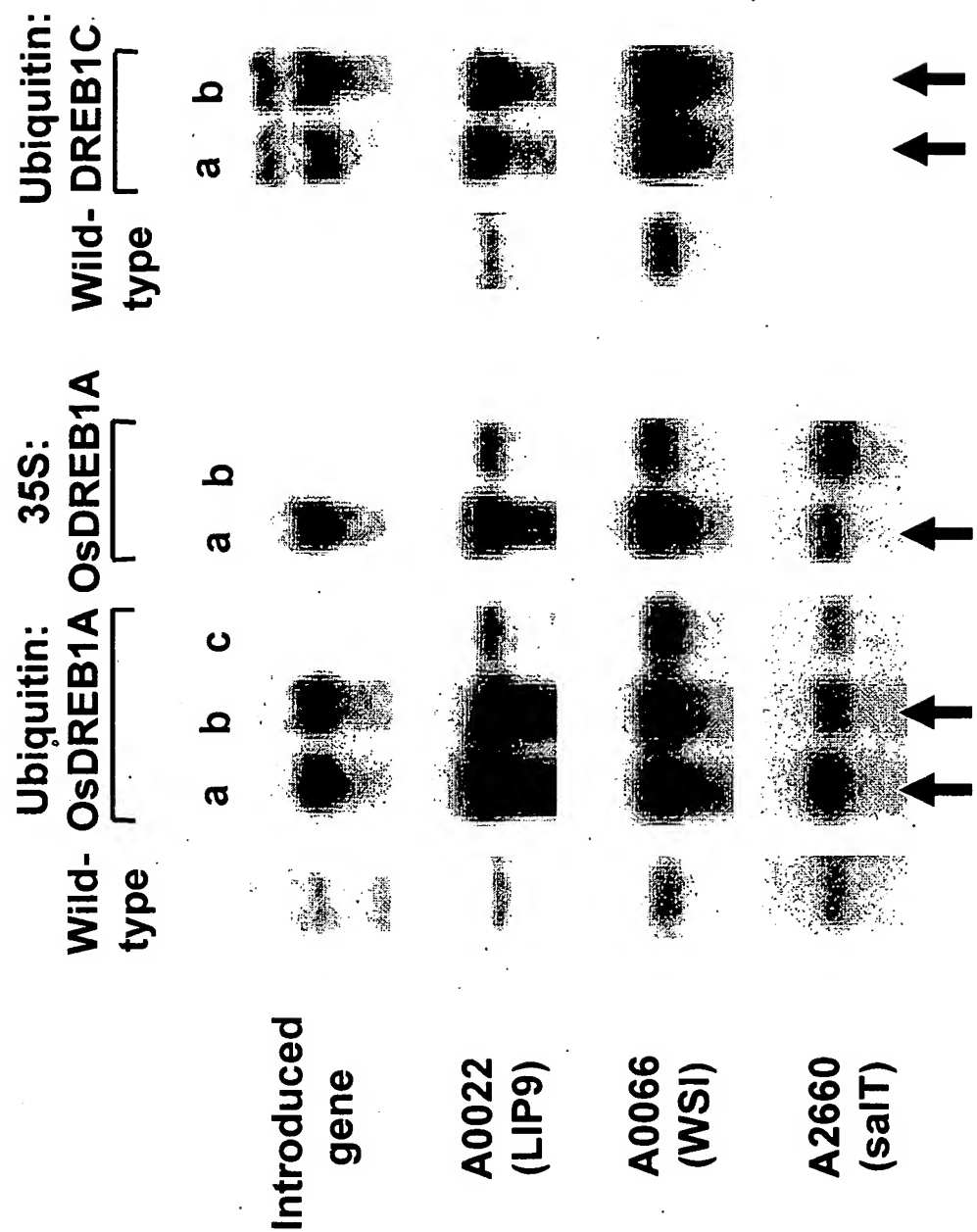


Fig. 6

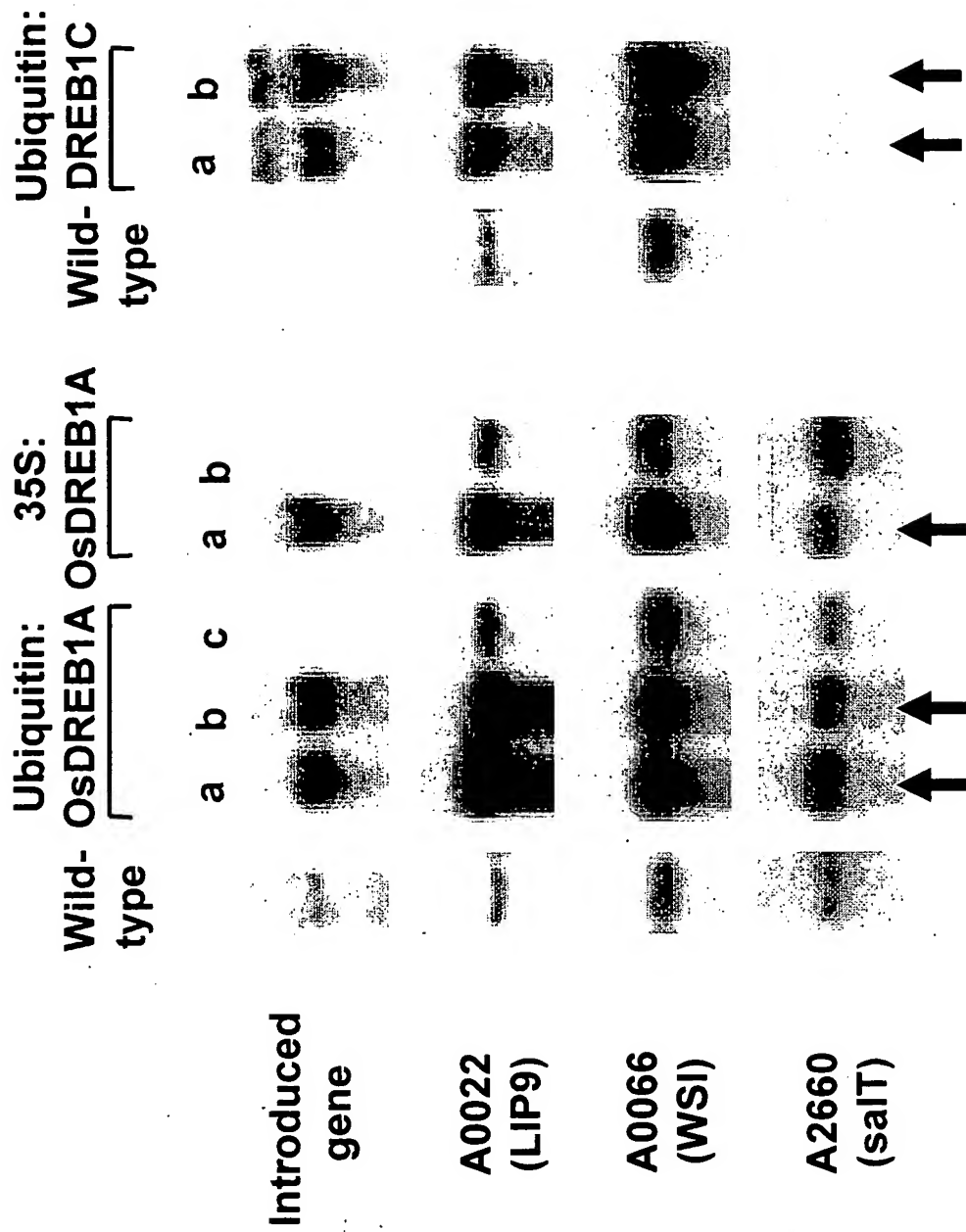


Fig. 7

WSI724

```

10      20      30      40      50      60
AGCCGTGGAAGTCCAACCTGCAGGCTCAGGCTGCAGATCGCCCAAGGCGCACTTGCCTCC

70      80      90      100     110     120
ACGATGGCTTGTCTCAACCGCTCGGAAGGCGAGATCCAATTGGCAATTTGTTCAACGCA

130     140     150     160     170     180
GGGAGAGAGGAGGAGACTGGAACGGGATCATTGGACATTGGTTGATGAATTGCAATTTGG

190     200     210     220     230     240
ATGACGAGGCCGCGAGGGTCAGACCGTCGGAGAGTGAGATGATGGTTATACAAGTGTACT

250     260     270     280     290     300
AGTAGGACGGACGGTGGCACCGGCCAGAAGCAGCAGATTTTGTGCAAACGTTGAGCCCCG

310     320     330     340     350     360
AACACGTGGCCGGCATCGACCCGCTACGACGGACGCAGCGCCCCCCCCCCCCCCCCCCCCC

370     380     390     400     410     420
GCGGACCCACGCGGGCCGGCCGCGCTGTCGCCGTGCTGCCGACTACGCCGTCGAAATCAA

430     440     450     460     470     480
CGCGTCCGCTCGATCCTCCCTGCCGACGCTGTACAAGTGGCGACCAAGAAACACCATG

490     500     510     520     530     540
TAGTATTTGATCTCGTCTAAGAGCAAGTTTAATACTATAGTCCACTATTAGCTCCAATTT

550     560     570     580     590     600
ATTTATAACTGATCTAATAGCCAATTCACACAATAATTGCTTACTATACTATTAATATAT

610     620     630     640     650     660
GGTCTCACATGTCATACACATATTCGGTCTTGGAGTTCTGCTGCAGCTGGCTACAGATC

670     680     690     700     710     720
TGTAGCCCGCTGCTCTTCTCTCAGAGCGAGTATAATAGTACAAACTGGACTGGCGATA

730     740     750     760     770     780
GGAGAAACACGTCAGCTACAGTGTGAGCTGGATGAGTGAGAAGAGGAGAGAGTGAGA

790     800     810     820     830     840
GTGGGCGACAATTTTATCGCCGGCTCTAGCACCAGCTTCGAGAGAAAAGTGGTGAGCGCA

850     860     870     880     890     900
GAGGTTGTGAGCTGCATGTGTGAGACGAAGCTTAAGTTATTTTATTATGATGTGAAGTTG

910     920     930     940     950     960
ATGGGTCCAGCGTTGCAGGTCATTTATTGTATTACAAAGATGCAAAGAGAGCTACTAGCT

970     980     990     1000    1010    1020
GAGTTGGATGGAATTAACGCCGGCTGTCTACGCTACTATTAACCTTGCTCTCATCTTTTA

1030    1040    1050    1060    1070    1080
TCTCATCAAAATATATTTATAGCTGGCTAATAGTCTGCTATCGTACCTGCTCTAATGCAT

1090    1100    1110    1120    1130    1140
ACGTTTTTTCTCTCTGTGGCAAACGGTTGGTGCGTTACACGGGGTGACGAAGCCATGC

1150    1160    1170    1180    1190    1200
ATCACCTGTCTCAACCGTCTCCTTTTTTAGCCTAATCTTTTCTCTTATCCGATGGGC

1210    1220    1230    1240    1250    1260
CTTCCGTTTCTCAAGACACCCCCACACCGCCCCGGCCCTCTATAAATACCAACCACGACG

1270    1280    1290    1300    1310    1320
AGCCAAGCGAACATCACCACAGCTAGATCATTAGCAATCCATTCCGATCCATCAAATTC

1330    1340    1350    1360
TCTTGAGACCGTAGAGAGAGAGAGAGGCGCCAACCATGGCCGGCATC

```

ABRE (370-380)  
DRE (430-440)  
DRE (510-520)  
TATA (1290-1300)  
FLcDNA (1340)  
ORF (1360)

WSI724



■ DRE(A/GCCGAC) ■ ABRE(ACGTGG/T) □ TATA box

■ cDNA

■ ORF

Fig. 8

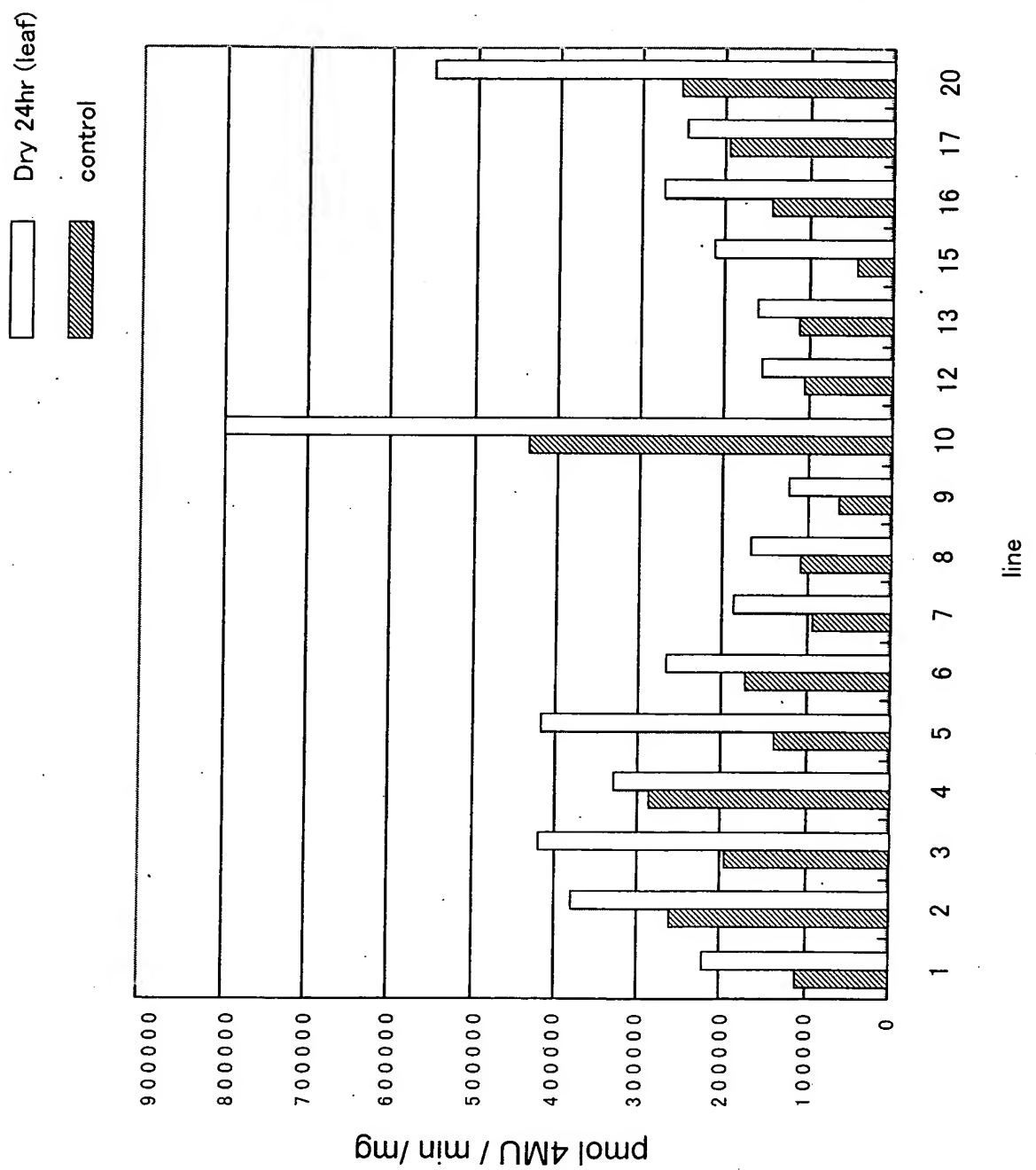


Fig. 9

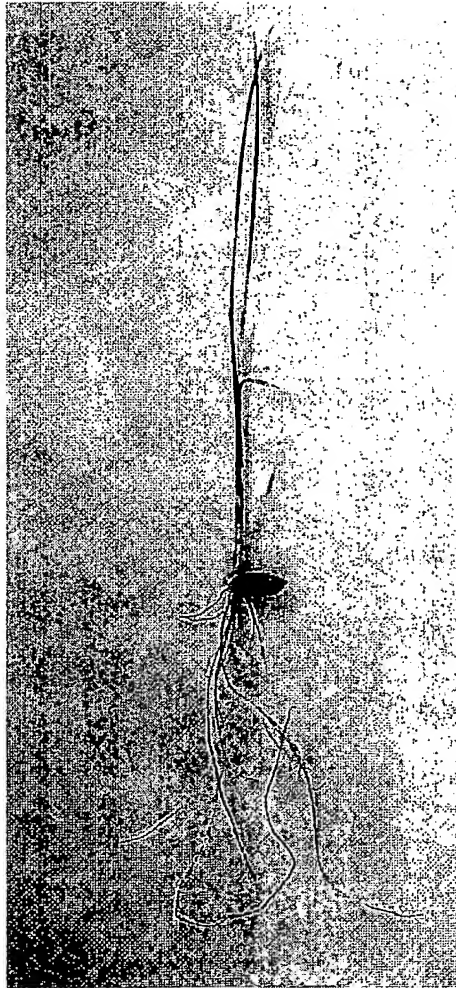


Fig. 9

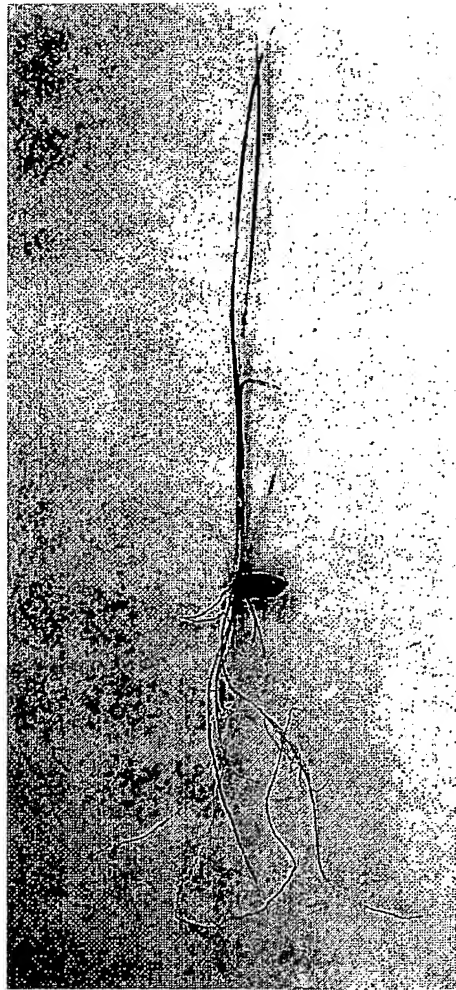


Fig. 9

